

<221> VARIANT
 <222> (9)...(9)
 <223> Xaa is Arg, Asn, Asp, Glu or Gly

<221> VARIANT
 <222> (10)...(10)
 <223> Xaa is Gln, Leu or Gly

<221> VARIANT
 <222> (11)...(11)
 <223> Xaa is Ala, Trp or Tyr

<221> VARIANT
 <222> (12)...(12)
 <223> Xaa is Ala, Gly, His, Phe, Thr or Val

<221> VARIANT
 <222> (14)...(14)
 <223> Xaa is Asn, Gln, Phe, Ser or Val

<221> VARIANT
 <222> (15)...(15)
 <223> Xaa is Arg, Leu, Pro or Ser

<221> VARIANT
 <222> (16)...(16)
 <223> Xaa is Leu, Ser, Trp or Tyr

<400> 1
 Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa
 1 5 10 15

<210> 2
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Family of preferred CEA binding moieties

<221> VARIANT
 <222> 1
 <223> Xaa is Asn or Asp

<221> VARIANT
 <222> 6
 <223> Xaa is Phe, Met, Leu or Asn

<221> VARIANT
 <222> 7
 <223> Xaa is Asp, Gly, Ile, Lys, Phe or Thr

<221> VARIANT
 <222> 9
 <223> Xaa is Arg, Asn, Asp, Glu, Gly or Trp

<221> VARIANT
<222> 12
<223> Xaa is Ala, Gly, His, Phe, Thr, Tyr or Val

<221> VARIANT
<222> (15)...(15)
<223> Xaa is Arg, Leu, Pro or Ser

<221> VARIANT
<222> (16)...(16)
<223> Xaa is Leu, Ser, Trp or Tyr

<400> 2
Xaa Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Xaa
1 5 10 15

<210> 3
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> CEA binding loop

<221> VARIANT
<222> 2
<223> Xaa is Asn, Glu or Met

<221> VARIANT
<222> 3
<223> Xaa is Asn, Leu, Met or Phe

<221> VARIANT
<222> 4
<223> Xaa is Asp, Gly, Ile, Lys, Phe or Thr

<221> VARIANT
<222> 5
<223> Xaa is Ala, Gln, Gly, Lys or Thr

<221> VARIANT
<222> 6
<223> Xaa is Arg, Asn, Asp, Glu or Gly

<221> VARIANT
<222> (7)...(7)
<223> Xaa is Gln, Gly or Leu

<221> VARIANT
<222> (8)...(8)
<223> Xaa is Ala, Trp or Tyr

<221> VARIANT
<222> (9)...(9)
<223> Xaa is Ala, Gly, His, Phe, Thr or Val

<400> 3

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
 1 5 10

<210> 4

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 4

Asn Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Ser Tyr
 1 5 10 15

<210> 5

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 5

Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Trp Thr Cys Asn Leu Leu
 1 5 10 15

<210> 6

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 6

Asn Trp Asp Cys Met Phe Gly Ala Glu Gly Trp Ala Cys Ser Pro Trp
 1 5 10 15

<210> 7

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 7

Asp Trp Val Cys Glu Lys Thr Thr Gly Gly Tyr Val Cys Gln Pro Leu
 1 5 10 15

<210> 8
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> CEA binding polypeptide

<400> 8
 Asn Trp Phe Cys Glu Met Ile Gly Arg Gln Trp Gly Cys Val Pro Ser
 1 5 10 15

<210> 9
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> CEA binding polypeptide

<400> 9
 Asp Trp Val Cys Asn Phe Asp Gln Gly Leu Ala His Cys Phe Pro Ser
 1 5 10 15

<210> 10
 <211> 12
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Parental domain for design of microprotein display library

<221> VARIANT
 <222> (1)...(12)
 <223> amino acid positions 4 and 9 are invariant Cys;
 all other positions Xaa are varied but not Cys, to
 provide a library of 2×10^8 different peptides
 based on the template sequence

<221> VARIANT
 <222> 1, 2, 3, 5, 6, 7, 8, 10, 11, 12
 <223> Xaa = Any Amino Acid except Cys

<400> 10
 Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa
 1 5 10

<210> 11
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Parental domain for design of microprotein display library

<221> VARIANT

<222> (1)...(11)

<223> amino acid positions 3 and 9 are invariant Cys; all other positions Xaa are varied but not Cys, to provide a library of 1×10^9 different peptides based on the template sequence

<221> VARIANT

<222> 1, 2, 4, 5, 6, 7, 8, 10, 11

<223> Xaa = Any Amino Acid except Cys

<400> 11

Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
1 5 10

<210> 12

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Parental domain for design of microprotein display library

<221> VARIANT

<222> (1)...(12)

<223> amino acid positions 3 and 10 are invariant Cys; all other positions Xaa are varied but not Cys, to provide a library of 1×10^9 different peptides based on the template sequence

<221> VARIANT

<222> 1, 2, 4, 5, 6, 7, 8, 9, 11, 12

<223> Xaa = Any Amino Acid except Cys

<400> 12

Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
1 5 10

<210> 13

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Parental domain for design of microprotein display library

<221> VARIANT

<222> (1)...(16)

<223> amino acid positions 4 and 13 are invariant Cys; all other positions Xaa are varied but not Cys, to

provide a library of 2.5×10^8 different peptides
based on the template sequence

<221> VARIANT

<222> 1, 2, 3, 5, 6, 7, 8, 9, 10, 11, 12, 14, 15, 16

<223> Xaa = Any Amino Acid except Cys

<400> 13

Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa
1 5 10 15

<210> 14

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Variable sublibrary sequence used in designing
focused secondary library

<221> VARIANT

<222> (1)...(3)

<223> Xaa is any amino acid except Cys

<221> VARIANT

<222> (5)...(6)

<223> Xaa is any amino acid except Cys

<221> VARIANT

<222> 1, 2, 3, 5, 6

<223> Xaa = Any Amino Acid except Cys

<400> 14

Xaa Xaa Xaa Cys Xaa Xaa Lys Lys Asp Gln Trp Thr Cys Asn Leu Leu
1 5 10 15

<210> 15

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Variable sublibrary sequence used in designing
focused secondary library

<221> VARIANT

<222> (5)...(9)

<223> Xaa is any amino acid except Cys

<221> VARIANT

<222> 5, 6, 7, 8, 9

<223> Xaa = Any Amino Acid except Cys

<400> 15
 Asp Trp Val Cys Xaa Xaa Xaa Xaa Xaa Gln Trp Thr Cys Asn Leu Leu
 1 5 10 15

<210> 16
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Variable sublibrary sequence used in designing
 focused secondary library

<221> VARIANT
 <222> (8)...(12)
 <223> Xaa is any amino acid except Cys

<221> VARIANT
 <222> 8, 9, 10, 11, 12
 <223> Xaa = Any Amino Acid except Cys

<400> 16
 Asp Trp Val Cys Glu Asn Lys Xaa Xaa Xaa Xaa Xaa Cys Asn Leu Leu
 1 5 10 15

<210> 17
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Variable sublibrary sequence used in designing
 focused secondary library

<221> VARIANT
 <222> (11)...(12)
 <223> Xaa is any amino acid except Cys

<221> VARIANT
 <222> (14)...(16)
 <223> Xaa is any amino acid except Cys

<221> VARIANT
 <222> 11, 12, 14, 15, 16
 <223> Xaa = Any Amino Acid except Cys

<400> 17
 Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Xaa Xaa Cys Xaa Xaa Xaa
 1 5 10 15

<210> 18
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Variable sublibrary sequence used in designing
 focused secondary library

 <221> VARIANT
 <222> (6)...(7)
 <223> Xaa is any amino acid except Cys

 <221> VARIANT
 <222> 9
 <223> Xaa is any amino acid except Cys

 <221> VARIANT
 <222> 12
 <223> Xaa is any amino acid except Cys

 <221> VARIANT
 <222> 15
 <223> Xaa is any amino acid except Cys

 <221> VARIANT
 <222> 6, 7, 9, 12, 15
 <223> Xaa = Any Amino Acid except Cys

 <400> 18
 Asp Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Leu
 1 5 10 15

 <210> 19
 <211> 16
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Variable sublibrary sequence used in designing
 focused secondary library

 <221> VARIANT
 <222> (5)...(7)
 <223> Xaa is any amino acid except Cys

 <221> VARIANT
 <222> 9
 <223> Xaa is any amino acid except Cys

 <221> VARIANT
 <222> 12
 <223> Xaa is any amino acid except Cys

 <221> VARIANT
 <222> 5, 6, 7, 9, 12
 <223> Xaa = Any Amino Acid except Cys

 <400> 19
 Asn Trp Val Cys Xaa Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Ser Tyr
 1 5 10 15

10/45

<210> 20
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Variable sublibrary sequence used in designing
 focused secondary library

<221> VARIANT
 <222> 1
 <223> Xaa is any amino acid except Cys

<221> VARIANT
 <222> 3
 <223> Xaa is any amino acid except Cys

<221> VARIANT
 <222> (14)...(16)
 <223> Xaa is any amino acid except Cys

<221> VARIANT
 <222> 1, 3, 14, 15, 16
 <223> Xaa = Any Amino Acid except Cys

<400> 20
 Xaa Trp Xaa Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Xaa Xaa Xaa
 1 5 10 15

<210> 21
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Isolate of TN10/9 library found not to bind CEA

<400> 21
 Asn Trp Arg Cys Lys Leu Phe Pro Arg Tyr Pro Tyr Cys Ser Ser Trp
 1 5 10 15

<210> 22
 <211> 15
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Isolate of TN10/9 library found not to bind CEA

<400> 22
 Arg Tyr Cys Glu Phe Phe Pro Trp Ser Leu His Cys Gly Arg Pro
 1 5 10 15

11/45

<210> 23
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Conserved amino acid positions in first family of
 CEA binding peptides

<221> VARIANT
 <222> 6
 <223> X is Asn, Leu, Met or Phe

<221> VARIANT
 <222> 7
 <223> X is Asp, Gly, Ile, Lys, Phe or Thr

<221> VARIANT
 <222> 9
 <223> X is Arg, Asn, Asp, Glu or Gly

<221> VARIANT
 <222> 12
 <223> X is Ala, Gly, His, Phe, Thr or Val

<221> VARIANT
 <222> 15
 <223> X is Arg, Leu, Pro or Ser

<400> 23
 Asp Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Leu
 1 5 10 15

<210> 24
 <211> 27
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetic CEA binding peptide with C-terminal
 immobilization sequence

<400> 24
 Ser Asn Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Ser
 1 5 10 15
 Tyr Ala Pro Gly Gly Glu Gly Gly Ser Lys
 20 25

<210> 25
 <211> 27
 <212> PRT
 <213> Artificial Sequence

12/45

<220>

<223> Synthetic CEA binding peptide with C-terminal immobilization sequence

<400> 25

Ser	Asp	Trp	Val	Cys	Glu	Asn	Lys	Lys	Asp	Gln	Trp	Thr	Cys	Asn	Leu
1				5					10					15	
Leu	Ala	Pro	Gly	Gly	Glu	Gly	Gly	Gly	Ser	Lys					
			20					25							

<210> 26

<211> 27

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic CEA binding peptide with C-terminal immobilization sequence

<400> 26

Ser	Asn	Trp	Asp	Cys	Met	Phe	Gly	Ala	Glu	Gly	Trp	Ala	Cys	Ser	Pro
1				5					10					15	
Trp	Ala	Pro	Gly	Gly	Glu	Gly	Gly	Gly	Ser	Lys					
			20					25							

<210> 27

<211> 27

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic CEA binding peptide with C-terminal immobilization sequence

<400> 27

Ser	Asp	Trp	Val	Cys	Glu	Leu	Thr	Thr	Gly	Gly	Tyr	Val	Cys	Gln	Pro
1				5					10					15	
Leu	Ala	Pro	Gly	Gly	Glu	Gly	Gly	Gly	Ser	Lys					
			20					25							

<210> 28

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> C-terminal sequence for immobilizing peptides

<400> 28

Ala	Pro	Gly	Gly	Glu	Gly	Gly	Gly	Ser	Lys
1				5				10	

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<210> 29
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Template sequence for sublibrary used in
 construction of focused secondary display library

<221> VARIANT
 <222> (1)...(3)
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> (5)...(6)
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> 1, 2, 3, 5, 6
 <223> Xaa = Any Amino Acid except Cys

<400> 29
 Xaa Xaa Xaa Cys Xaa Xaa Lys Lys Asp Gln Trp Thr Cys Asn Leu Leu
 1 5 10 15

<210> 30
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Template sequence for sublibrary used in
 construction of focused secondary display library

<221> VARIANT
 <222> (5)...(9)
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> 5, 6, 7, 8, 9
 <223> Xaa = Any Amino Acid except Cys

<400> 30
 Asp Trp Val Cys Xaa Xaa Xaa Xaa Xaa Gln Trp Thr Cys Asn Leu Leu
 1 5 10 15

<210> 31
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Template sequence for sublibrary used in
 construction of focused secondary display library

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<221> VARIANT
 <222> (8)...(12)
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> 8, 9, 10, 11, 12
 <223> Xaa = Any Amino Acid except Cys

<400> 31
 Asp Trp Val Cys Glu Asn Lys Xaa Xaa Xaa Xaa Xaa Cys Asn Leu Leu
 1 5 10 15

<210> 32
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Template sequence for sublibrary used in
 construction of focused secondary display library

<221> VARIANT
 <222> (11)...(12)
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> (14)...(16)
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> 11, 12, 14, 15, 16
 <223> Xaa = Any Amino Acid except Cys

<400> 32
 Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Xaa Xaa Cys Xaa Xaa Xaa
 1 5 10 15

<210> 33
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Template sequence for sublibrary used in
 construction of focused secondary display library

<221> VARIANT
 <222> (6)...(7)
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> 9
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> 12
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> 15
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> 6, 7, 9, 12, 15
 <223> Xaa = Any Amino Acid except Cys

<400> 33
 Asp Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Leu
 1 5 10 15

<210> 34
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Template sequence for sublibrary used in
 construction of focused secondary display library

<221> VARIANT
 <222> (5)...(7)
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> 9
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> 12
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> 5, 6, 7, 9, 12
 <223> Xaa = Any Amino Acid except Cys

<400> 34
 Asn Trp Val Cys Xaa Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Ser Tyr
 1 5 10 15

<210> 35
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Template sequence for sublibrary used in
 construction of focused secondary display library

<221> VARIANT
 <222> 1
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> 3
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> (14)...(16)
 <223> X is any amino acid except Cys

<221> VARIANT
 <222> 1, 3, 14, 15, 16
 <223> Xaa = Any Amino Acid except Cys

<400> 35
 Xaa Trp Xaa Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Xaa Xaa Xaa
 1 5 10 15

<210> 36
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Family of CEA binding polypeptides

<221> VARIANT
 <222> 1
 <223> Xaa is Asp, Asn, Ala or Ile

<221> VARIANT
 <222> 3
 <223> Xaa is Val, Ile, Met, Tyr, Phe, Pro or Asp

<221> VARIANT
 <222> 5
 <223> Xaa is Asn, Glu or Asp

<221> VARIANT
 <222> 6
 <223> Xaa is Leu, Phe, Tyr, Trp, Val, Met, Ile or Asn

<221> VARIANT
 <222> 7
 <223> Xaa is Phe, Leu, Asp, Glu, Ala, Ile, Lys, Asn,
 Ser, Val, Trp or Tyr

<221> VARIANT
 <222> (8)...(8)
 <223> Xaa is Lys, Phe, Asp, Gly, Leu, Asn or Trp

<221> VARIANT
 <222> (9)...(9)
 <223> Xaa is Asn, Pro, Phe, Gly, Asp, Ala, Ser, Glu, Gln

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or Trp

<221> VARIANT

<222> (10)...(10)

<223> Xaa is Gln or Lys

<221> VARIANT

<222> (12)...(12)

<223> Xaa is Phe, Thr, Met, Ser, Ala, Asn, Val, His,
Ile, Pro, Trp or Tyr

<221> VARIANT

<222> (14)...(14)

<223> Xaa is Asn, Asp, Glu, Pro, Gln or Ser

<221> VARIANT

<222> (15)...(15)

<223> Xaa is Val, Leu, Ile, Pro, Ala, Gln, Ser, Met,
Glu, Thr, Lys or Trp

<221> VARIANT

<222> (16)...(16)

<223> Xaa is Leu, Met, Val, Tyr, Ala, Ile, Trp, His,
Pro, Gln, Glu, Phe, Lys or Arg

<400> 36

Xaa	Trp	Xaa	Cys	Xaa	Xaa	Xaa	Xaa	Xaa	Trp	Xaa	Cys	Xaa	Xaa	Xaa
1			5					10				15		

<210> 37

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 37

Asp	Trp	Met	Cys	Asn	Leu	Phe	Lys	Asn	Gln	Trp	Phe	Cys	Asp	Leu	Met
1				5					10					15	

<210> 38

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 38

Asp	Trp	Val	Cys	Asn	Leu	Phe	Lys	Asn	Gln	Trp	Phe	Cys	Asp	Leu	Met
1				5					10					15	

<210> 39
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> CEA binding polypeptide

<400> 39
 Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Gln Met
 1 5 10 15

<210> 40
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> CEA binding polypeptide

<400> 40
 Asn Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Gln Glu
 1 5 10 15

<210> 41
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> CEA binding polypeptide

<400> 41
 Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Gln Val Lys
 1 5 10 15

<210> 42
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> CEA binding polypeptide

<400> 42
 Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Met
 1 5 10 15

<210> 43
 <211> 16
 <212> PRT
 <213> Artificial Sequence

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<220>

<223> CEA binding polypeptide

<400> 43

Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Gln Ile
1 5 10 15

<210> 44

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 44

Ile Trp Asp Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Pro Ala Pro
1 5 10 15

<210> 45

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 45

Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Ile Arg
1 5 10 15

<210> 46

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 46

Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Val
1 5 10 15

<210> 47

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 47

Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Ala Ile
1 5 10 15

<210> 48

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 48

Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Met Ala
1 5 10 15

<210> 49

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 49

Asp Trp Val Cys Glu Phe Leu Lys Met Gln Trp Ala Cys Asn Val Leu
1 5 10 15

<210> 50

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 50

Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Val Met
1 5 10 15

<210> 51

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> CEA binding polypeptide

<400> 51

Ala Trp Pro Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Pro Pro Gln
1 5 10 15

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<210> 52
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> CEA binding polypeptide

<400> 52
 Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Leu
 1 5 10 15

<210> 53
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> CEA binding polypeptide

<400> 53
 Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Lys Trp
 1 5 10 15

<210> 54
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> CEA binding polypeptide

<400> 54
 Asp Trp Val Cys Glu Trp Leu Lys Met Gln Trp Ala Cys Asn Met Leu
 1 5 10 15

<210> 55
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> CEA binding polypeptide

<400> 55
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33/45

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1 5 10 15

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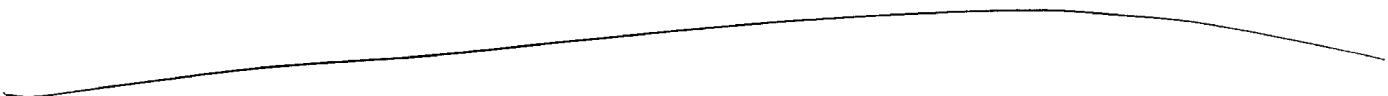
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A DOCPHOENIX

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Specification Not in English

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____ N/AP _____
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____ REM _____
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____ SRNT _____
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____ CLMPTO _____
PTO Prepared Complete Claim Set

____ ECBOX _____
Evidence Copy Box Identification

____ WCLM _____
Claim Worksheet

____ WFEE _____
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File Wrapper

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